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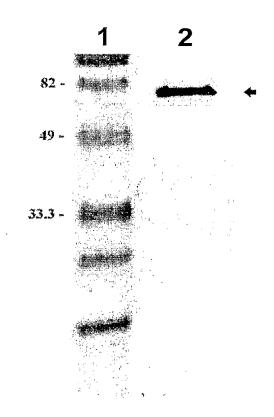


FIGURE 1

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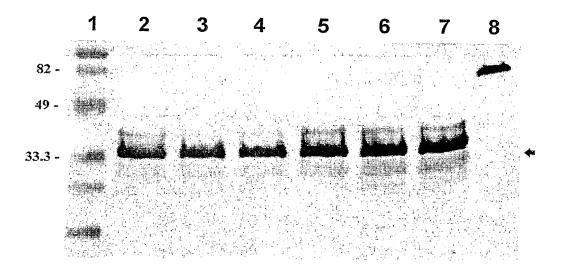


FIGURE 2

-101	AGCGTTTAGC	TGAACTTGAA	GCTATGGTAA	AAGAGCTTGA	<u>AA</u> CTTTAGAG	
				-35		
-51	CAAGAGCAAC	AAAGCTAATA	TTT <u>TAATAT</u> T	TT.AAGGAGTA	TAAAAGAAAA	
			-10	SD		
1	<b>ATG</b> AGTACTT	TAATAAAGAA	AATCGTAGCT	TATATAGCTT	TAATCTCTTT	
51	TAGTTTTAGC	GTATTACCTG	CTCAAACTTA	TGATGATGCG	GCTAGAATTA	
101	CTGGAGAAGC	TGAGACTTTA	CAAAATGACG	GAGAATACCA	AAAGTCTTAT	
151	GATAAATCTC	AAGAGGCTTC	TGACTCTATA	GATAAAACTA	CTGTATCATT	
201	ATTTTATAGA	TTAATGAACT	TAAGAATAGC	TAAAGCAAAA	AATGATGCAA	
251	ATAAGACTAT	TAATGAAATA	GAACAATTAG	GTGCTTCTAC	TGATAATGAA	
301	TTTAAAACAA	AATATCAAGA	AGCTCTAAAA	TTCTTTGAAG	AAGGAAATAA	
351	TAGTATTACT	AACTTACCTC	CAGAACCGCA	AACTCCTCCT	ACAGATGAAG	
401	AGTTTACTGC	TTCTTCAAAC	ACATTCACTA	CAGTATATAA	TTCTTTCAAC	
451	AATGCTTTAC	AATCTGCTAA	CAGTGTAAAA	GAAGGTTATC	TTAATAGAGA	
501	AAGAGCAATA	GCTTCAAAAT	CCATTAATGA	TGCTAGAAAC	AAATATAAAG	
551	CAGAATTAGG	CAAGAGTGTA	AAAGCAGGCG	ATGCTAATGA	TAGAAATATT	
601	AATGGTGCTT	TAACTAGAGC	TGATGAAGCA	CTTAGCAATG	ACAATTTTGC	
651	AAGCGTTCAG	CAGAATGTAT	CTACTGCATT	AGCTGGTATA	AATAAAGCTA	
701	TAGCAGATGC	TAAGGCGAAA	GCTGAGGCAG	AAGCTAAAGC	AAAAGCTGCT	
751	GCTGAAGCTA	AGGCAAGAGC	TGAAGCAGAG	GCTAAAGCGA	AAGCAGAAGC	
801	TGCTGCTAAA	GCAAAAGCTG	AAGCAGAGGC	TAAAGCGAAA	GCAGATGCAA	
851	TAGCAAAAGC	TAAAAAAGAC	ATAGAAGATG	CACAAAATAA	ATATAATAAT	
901	TTAGTTAATG	ATCAAGTAAT	AGCTAAAGGT	GATGATAATG	ATAAAAACGT	
951	ATCAAAACTT	TTAACTGATG	CTAATAATGC	TTTACAAAAC	ACTCCTCAAA	
1001	CTGCAAGCGA	TAAAGCTTTA	GAAGCTTCTA	AAACTATGGA	TAATATATTA	
1051	AACACTGCTA	ATCAATTGAA	AAAAGAAGAA	GCTGTTAAAA	ATCTAGAGCA	
1101	ATTAAAGGCA	AGAAGAGACA	GACTTATAAG	CGAAGGTTAT	TTAACTAAAG	
1151	ACAGCGAAGA	AGAACAAAAG	TTATCTCAAA	CTATTAAAGA	AGCTGAAGAT	
1201	GCTTTAAATA	ACAATGATTA	TGTTTTAGCT	GACCAAAAAA	TGCAGGAAGC	
1251	TAATCTTAAC	ATGAATGCTA	TAGAAGAGAG	AGGACCTATT	GACGGACAAG	
1301	TTATACCTGG	TGAAATGGGC	GGTAACGAAA	CTGGTCAAAT	AATTGATGCT	
1351	ACTACTGGTC	AAGAAGTAAA	TACAGAAGGA	AAAGTTACTG	TATTACCTCA	
1401	ATATTATGTT	GTAGTAAGAA	GAGTACCTCT	AACTGATGCT	TTATGGAGAA	
1451	TTGCTGGATA	CAGCTACATA	TACAACAACC	CTATAGAATG	GTACAGAATA	
1501	TATGAAGCTA	ACAGAAATGT	ACTTAGAGAC	CCTAATAACC	CTGATTTAAT	
1551	ACTTCCTGGT	CAAAGATTAA	TAATACCTAG	CCTTAATGGT	GAAGAGAGAA	
1601	GCGGTGATTA	TAATCCTGAT	TTAGAGTATT	TGACTTATGA	TGAGGTTATG	
1651	CAGTTAAGAC	AGCAAAATAA	CACTACTCAA	GCACAACAGT	<b>AA</b> GAAATAAA	
1701	CTTATAAAAT	ACAAAAGGTC	ATGCATTTAA	TATGTATGAC	CTTTTTTTGT	

## FIGURE 3

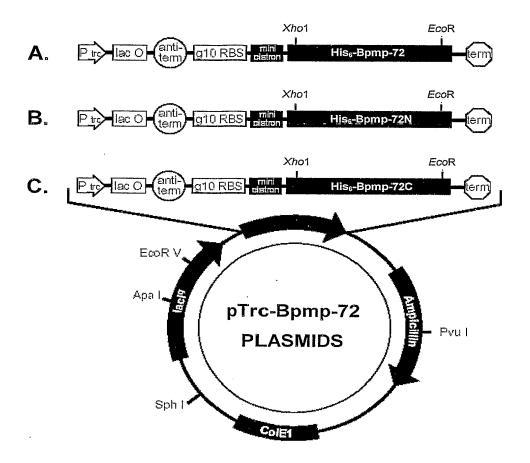


FIGURE 4

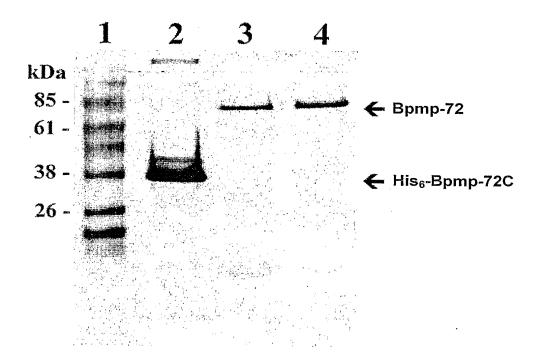


FIGURE 5

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FIGURE 6

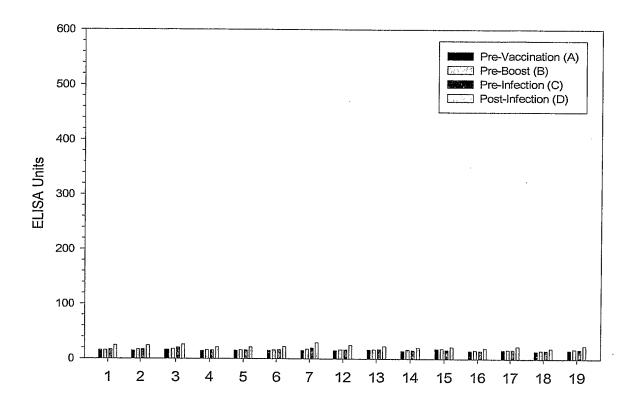


FIGURE 7

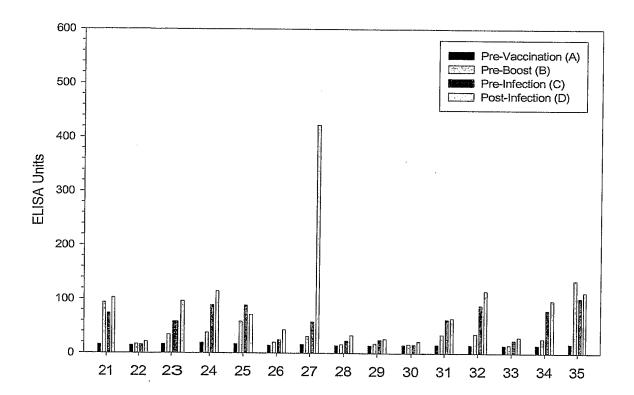


FIGURE 8

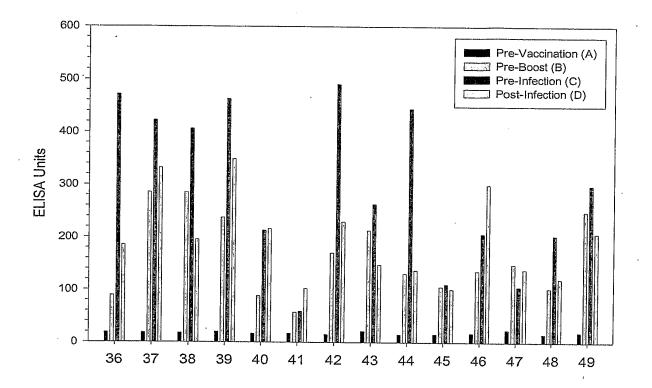


FIGURE 9

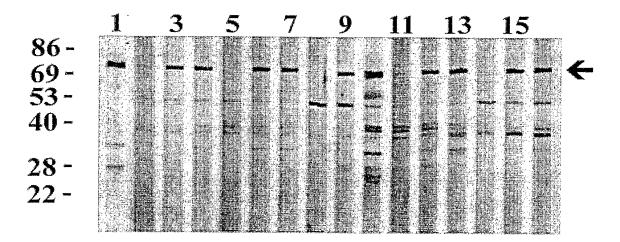


FIGURE 10

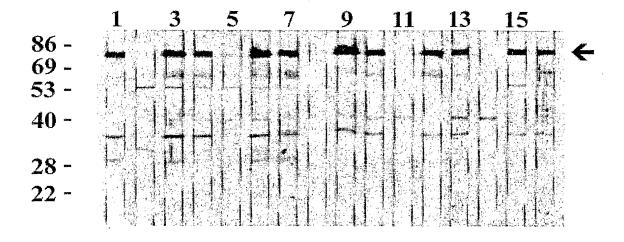
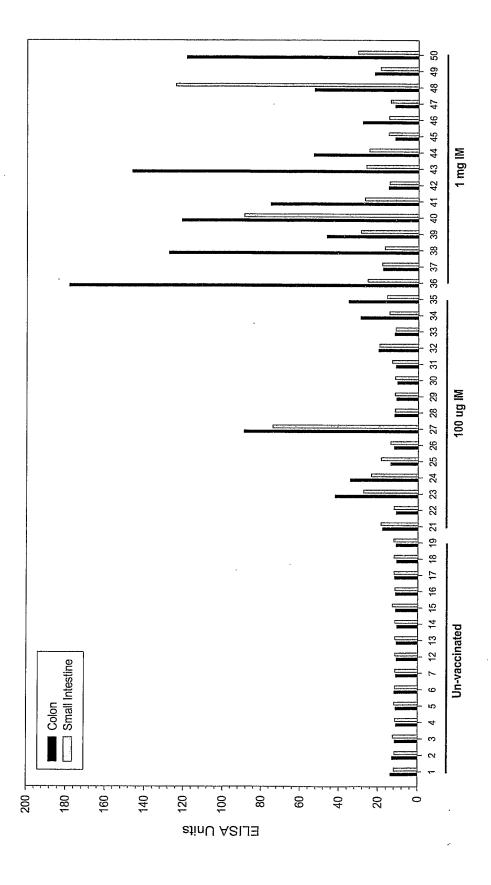


FIGURE 11



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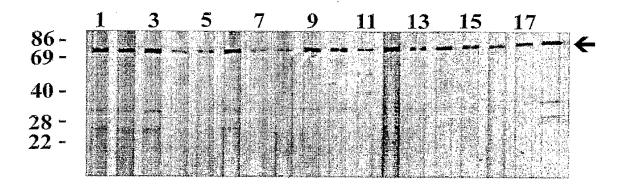


FIGURE 13